

Finding the Needle...Again...and Again...and Again: Lessons Learned from NCI Programs

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Health Sciences Director

Deputy Director, Center for Strategic Scientific Initiatives (CSSI)

Deputy Director, Center for Cancer Genomics (CCG)

Office of the Director, National Cancer Institute (NCI)

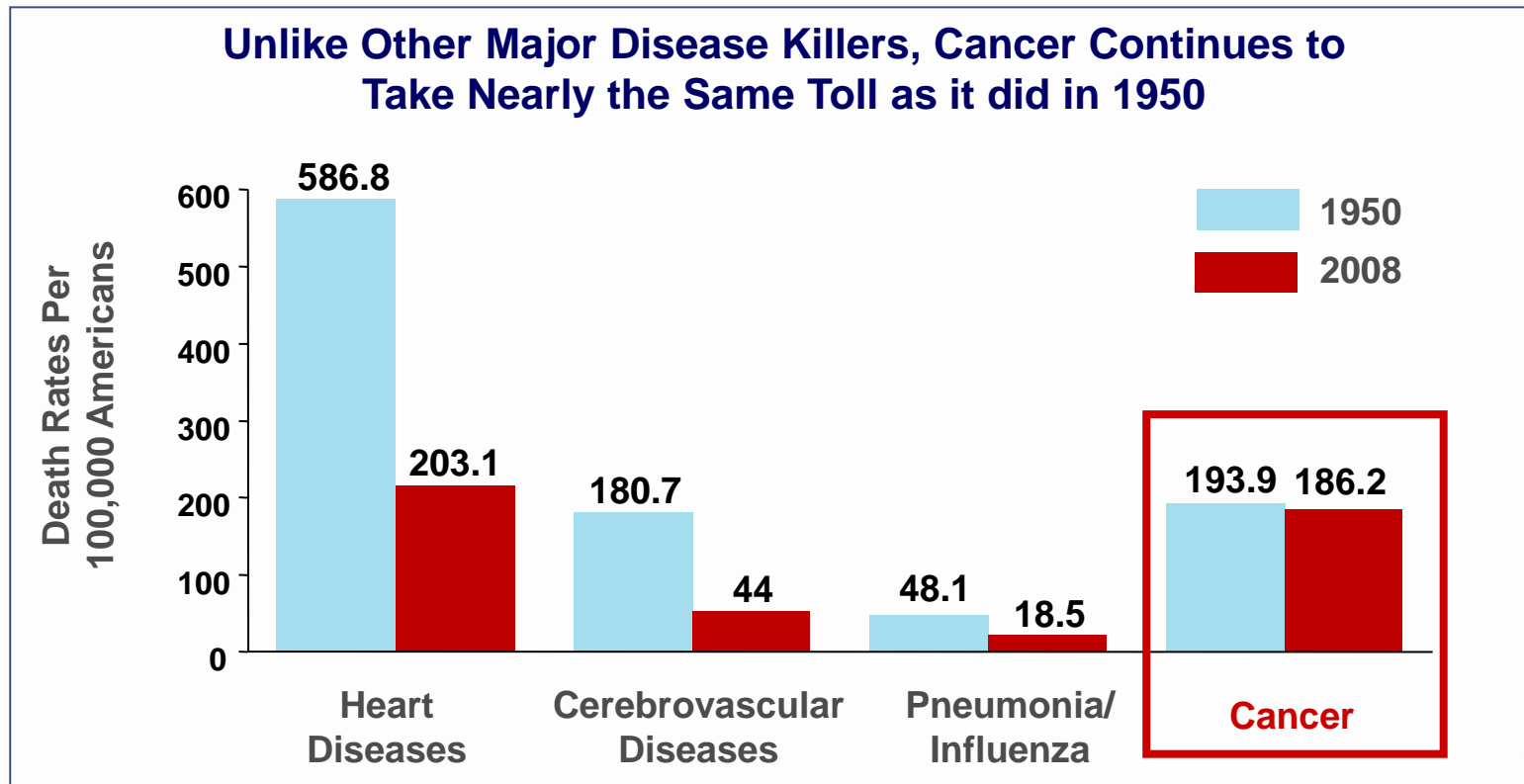
National Institutes of Health (NIH)

World CTC Summit, Boston MA

November 13, 2012

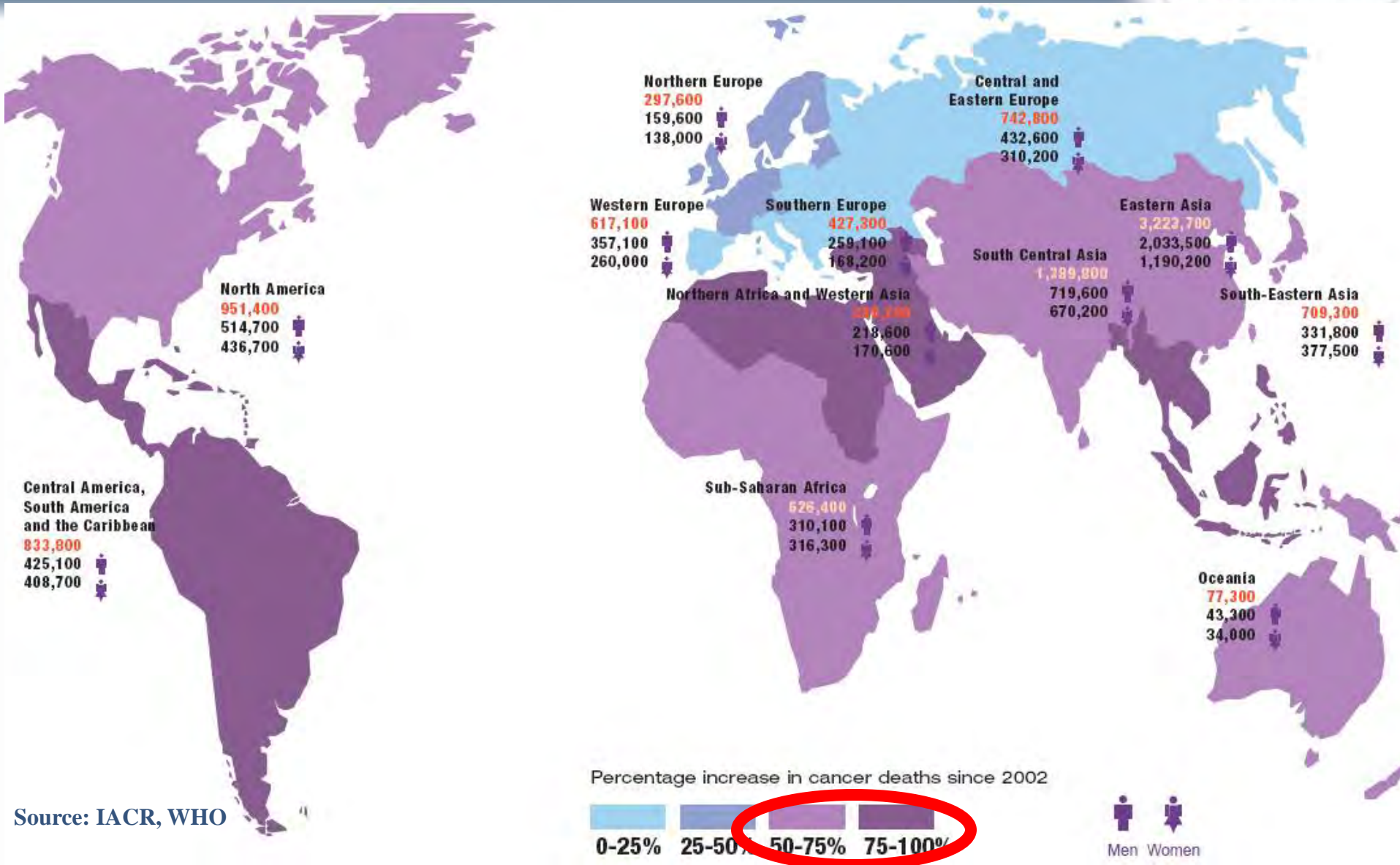
In the U.S., Cancer Continues to Represent an Enormous Burden

- **569,490** Americans died of cancer in 2010
- **1,529,560** Americans will be diagnosed with cancer this year
- **\$124.6 billion** in 2010 for cancer healthcare costs



Source for 2010 deaths and diagnoses: American Cancer Society (ACS) 2010 Cancer Facts & Figures; Atlanta, Georgia
Source for 2008 age-adjusted death rate: National Center for Health Statistics, NCHS Public-use file for 2008 deaths.

Global Burden: By 2020, Cancer Mortality 10 M/yr (Incidence 16 M/yr)



Unprecedented Amount of Scientific Knowledge: Omics(ssss)

A map of human genome variation from
population-scale sequencing

The 1000 Genomes Project Consortium



2001



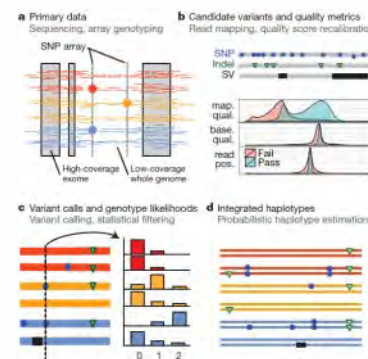
2010

NATURE

1 NOVEMBER 2012

An integrated map of genetic variation
from 1,092 human genomes

The 1000 Genomes Project Consortium



2012

2005

1923

49,024 pubs

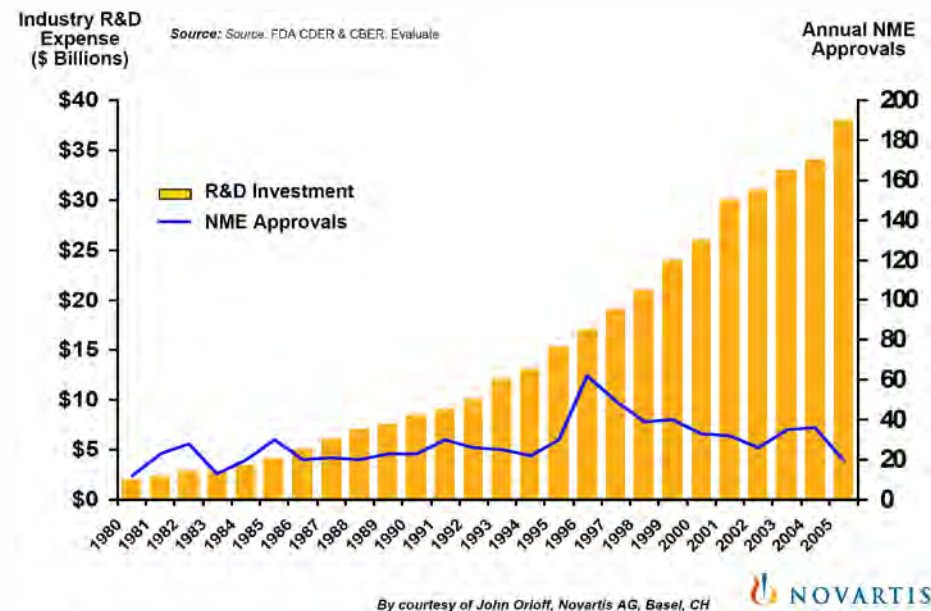
54,587 pubs

87,793 pubs

38,506 pubs

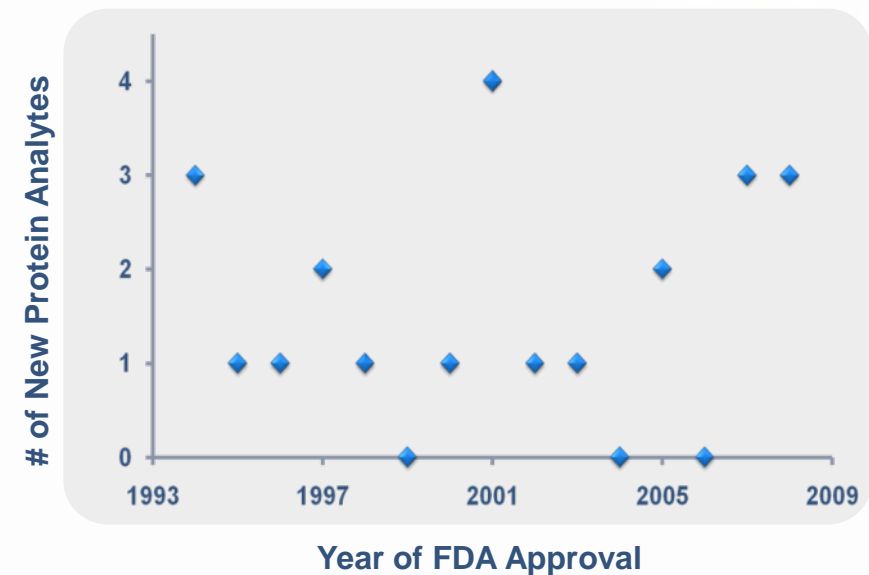
Is More Knowledge Yielding More Solutions for Patients?

Drug Discovery and Development



- 10 – 15 years at ~ \$1.8 billion*
- 2007: 19 NMEs [lowest since 1983]
- 2008: 21 NMEs [29% new-in-class]
- 2009: 24 NMEs [17% new-in-class]

Diagnostic Biomarkers



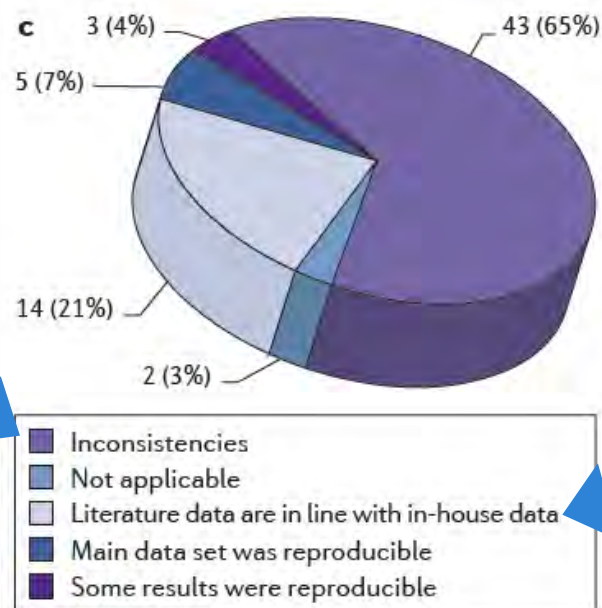
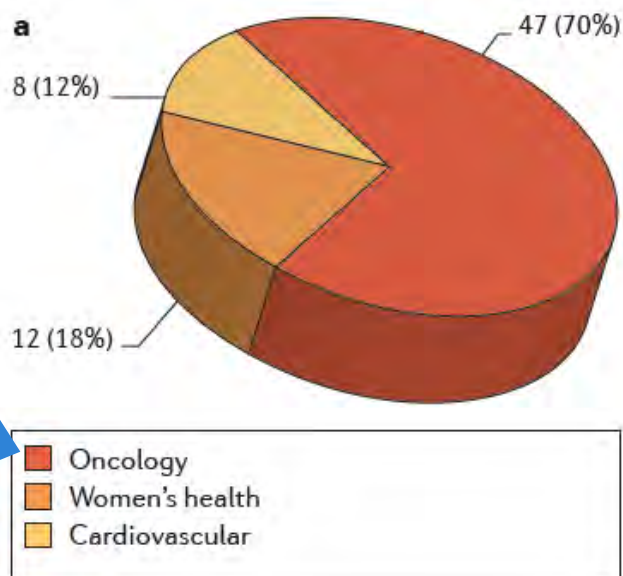
- Averaging 1.5 FDA approvals per year†
- 1000's of samples
- Balancing complexity of biology against heterogeneity of patients

Maybe...but can it be more efficient?

Too Many Papers: Is It Just Overwhelming Literature?

NATURE REVIEWS | DRUG DISCOVERY

Believe it or not: how much can we rely on published data on potential drug targets?



Innovation Exists in Public Sector Research Institutions (PSRIs)

Table 1. Number of Drug Products Approved by the Food and Drug Administration and Originating from Public-Sector Research, According to Therapeutic Area, 1970–2009.

Therapeutic Area	Number
Total	153
Hematology or oncology	40
Infectious disease	36
Cardiology	12
Metabolic disease	12
Central nervous system	12
Dermatology	7
Renal disease	7
Ophthalmology	6
Immunology	6
Gastroenterology	4
Women's health	3
Allergy	2
Pulmonary disease	2
Urology	2
Anesthesiology	1
Dental disorders	1

Over the past 40 years,

- **153** FDA approvals were carried out in PSRIs (~9.3% of overall approvals)

- 93 small molecule
- 36 biologics
- 15 vaccines
- 8 IVDs
- 1 over-the-counter drug

- Most prolific PSRIs include:

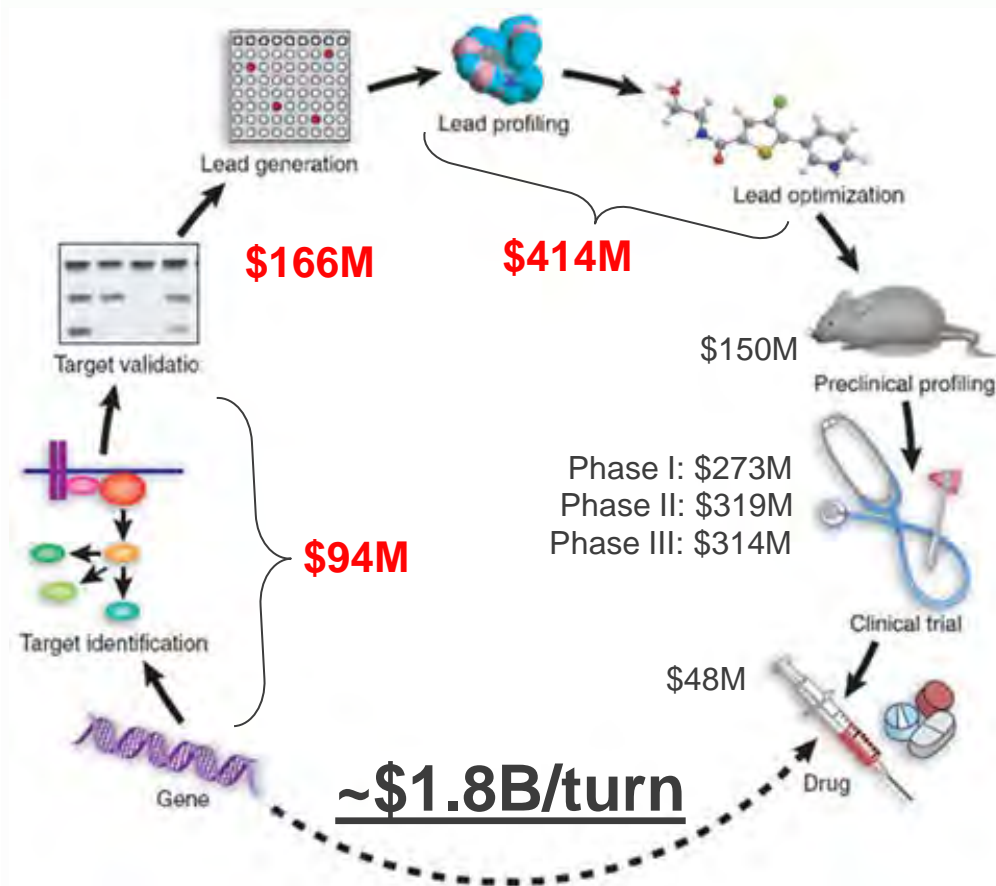
- NIH (22 products)
- UC system (11 products)
- MSKCC (8 products)
- Emory (7 products)
- Yale (6 products)

- Virtually all important, innovative vaccines that have been introduced in the last 25 years have been created by PSRIs

Table 2. FDA-Approved Drugs Discovered through Public-Sector Research, According to Type of Review and Chemical Type, 1990–2007.*

Type of Review	New Molecular Entity	New Ester, Salt, or Derivative	New Formulation	New Combination	New Manufacturer	New Indication	Already Marketed	Total
Priority review								
Discovered by PSRI (no.)	44	1	17	3	0	1†	0	66
All FDA approvals (no.)	209	6	99	20	14	0	0	348
Rate of PSRI discovery (%)	21.1	16.7	17.2	15.0	0	NA	NA	19.0
Standard review								
Discovered by PSRI (no.)	20	0	36	6	7	8	0	77
All FDA approvals (no.)	274	33	631	96	137	10	12	1193
Rate of PSRI discovery (%)	7.3	0	5.7	6.3	5.1	80.0	0	6.5
All approvals								
Discovered by PSRI (no.)	64	1	53	9	7	9	0	143
All FDA approvals (no.)	483	39	730	116	151	10	12	1541
Rate of PSRI discovery (%)	13.3	2.6	7.3	7.8	4.6	90.0	0	9.3

Translation Pace: How To Break Out of Current Paradigm?



Turning the Crank...

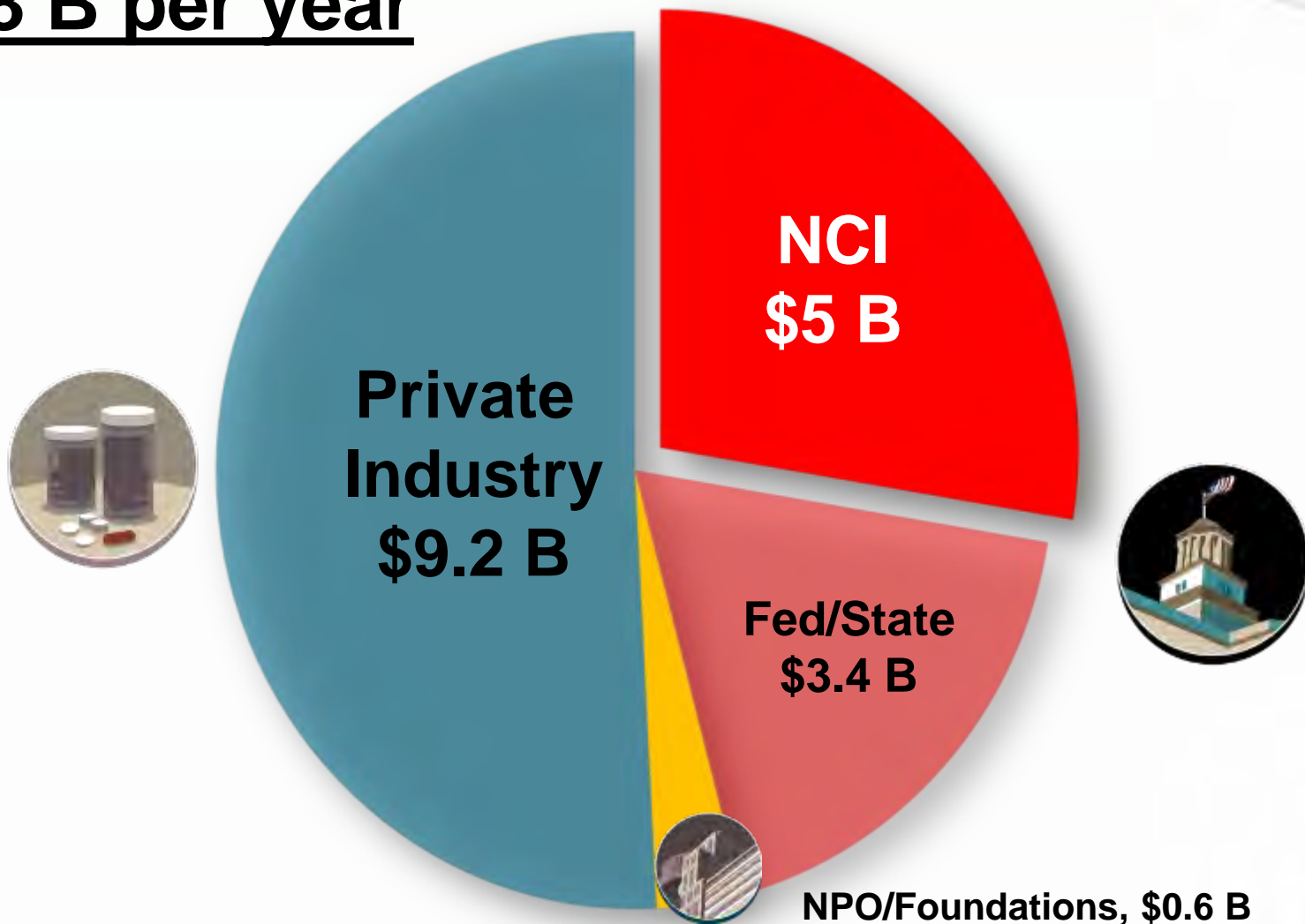
Key Needs (from community '02)

- Standards and protocols
- Real-time, public release of data
- Large, multi-disciplinary teams
- Pilot-friendly team environment to share failures and successes
- Team members with **trans-disciplinary training**

The potential to transform cancer drug discovery and diagnostics

National Cancer Program: Stakeholders

~\$18 B per year



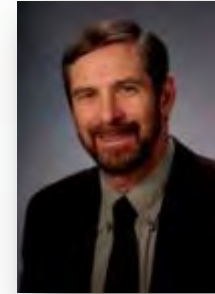
National Cancer Institute Organization



Director
Harold Varmus, MD

**National Cancer
Institute**

\$5.07B
(FY12)



Deputy Director
Douglas Lowy, MD

**Office of the
Director**

CSSI

CCG

~\$190 M (~4%)

**Center for
Cancer
Research**

**Division of
Cancer
Epidemiology
and Genetics**

**Division of
Cancer
Treatment
and
Diagnosis**

**Division of
Cancer
Biology**

**Division of
Cancer
Control and
Population
Sciences**

**Division of
Cancer
Prevention**

**Division of
Extramural
Activities**

Conducting – Intramural

Funding – Extramural

NCI Center for Strategic Scientific Initiatives (CSSI): Concept Shop



Director
Douglas Lowy, MD



~\$190M (FY12)



Deputy Director
Jerry S.H. Lee, PhD



**Office of Cancer Clinical
Proteomics Research**

Director
Henry Rodriguez, PhD, MBA



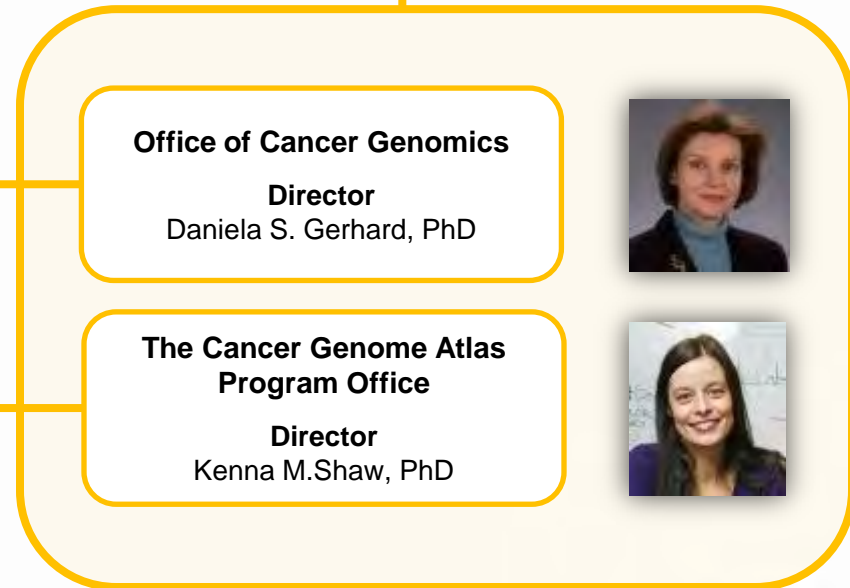
**Office of Physical Sciences-
Oncology**

Director
Larry A. Nagahara, PhD



**Office of Cancer
Nanotechnology Research**

Director
Piotr Grodzinski, PhD



Office of Cancer Genomics

Director
Daniela S. Gerhard, PhD



**The Cancer Genome Atlas
Program Office**

Director
Kenna M. Shaw, PhD



*Center for Cancer Genomics (CCG) shown in yellow

NCI Center for Strategic Scientific Initiatives (CSSI): Concept Shop



Director
Douglas Lowy, MD



~\$190M (FY12)



Deputy Director
Jerry S.H. Lee, PhD

Mission

“...to create and uniquely implement exploratory programs focused on the development and integration of advanced technologies, trans-disciplinary approaches, infrastructures, and standards, to accelerate the creation and broad deployment of data, knowledge, and tools to empower the entire cancer research continuum in better understanding and leveraging knowledge of the cancer biology space for patient benefit...”



2003, 2007, 2011



2005, 2010



2008



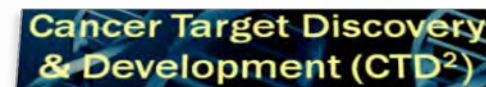
2011



2004, 2008



2005, 2008

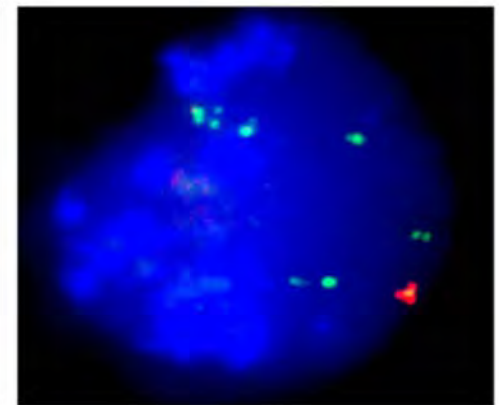
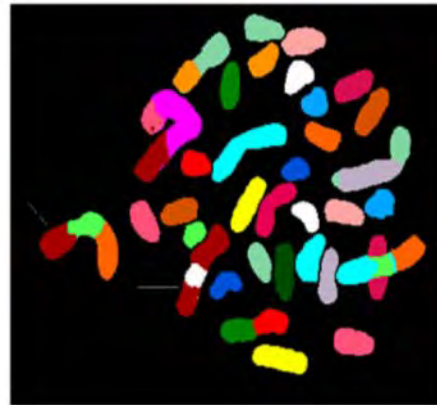


2010

Example #1: Cancer Genomics- Taking a Page from Engineers

Disease of Genomic Alterations

- Copy number
- Expression (regulation of)
- Regulation of translation
- Mutations
- Epigenome



- **Systematic identification of all genomic changes**
- **Repeat (a lot) for individual cancer**
- **Repeat for many cancers**
- **Make it publically available**

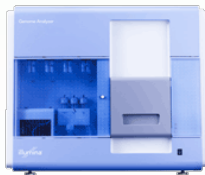
Saturated steam				Superheated steam		
Pressure (kg/cm ²)	Temp (°C)	Vapour enthalpy (kcal/kg)	Specific volume (m ³ /kg)	Density (kg/m ³)	Specific volume (m ³ /kg) at 250°C at 300°C	
1	99.1	638.8	1.725	0.580	2.454	2.691
2	119.6	646.2	0.902	1.109	1.223	1.342
3	132.9	650.6	0.617	1.621	0.812	0.893
4	142.9	653.7	0.471	2.123	0.607	0.668
5	151.1	656.0	0.382	2.618	0.484	0.533
6	158.1	657.0	0.321	3.115	0.402	0.443
7	164.2	659.5	0.278	3.597	0.343	0.379
8	169.6	660.8	0.245	4.082	0.299	0.331
9	174.5	661.9	0.219	4.566	0.265	0.293
10	179.1	662.9	0.198	5.051	0.238	0.263
12	187.1	664.5	0.166	6.024	0.196	0.218
14	194.1	665.7	0.143	6.993	0.167	0.186
16	200.4	666.7	0.126	7.937	0.145	0.162
18	206.1	667.4	0.112	8.929	0.128	0.143
20	211.4	668.0	0.101	9.901	0.114	0.128
22	216.2	668.4	0.092	10.870	0.103	0.116
24	220.7	668.7	0.085	11.765	0.093	0.106
26	225.0	669.0	0.078	12.821	0.085	0.097
28	229.0	669.1	0.073	13.699	0.078	0.089
30	232.7	669.2	0.068	14.706	0.072	0.083

Steam table (Reference)

Many “Thermometers”: Heterogeneity of Platforms



454



Illumina



SOLiD

Complete Genomics
Complete Genomics



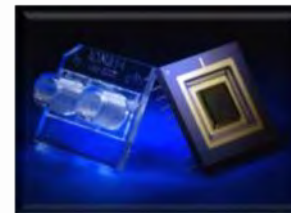
Helicos



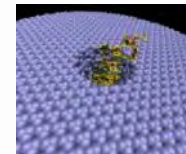
Visigen



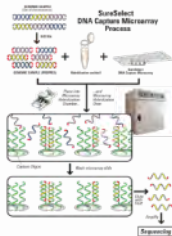
PacBio



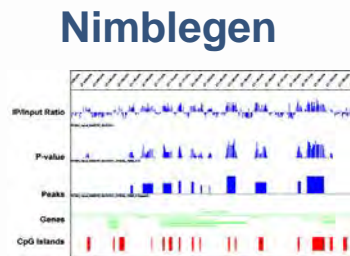
Ion-Torrent



Oxford Molecular



Agilent

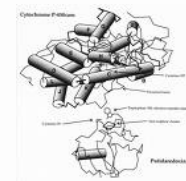


Nimblegen



LaserGen

ZSGenetics



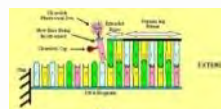
NABsys



Raindance



Febit



Intelligent Biosystems



Halycon

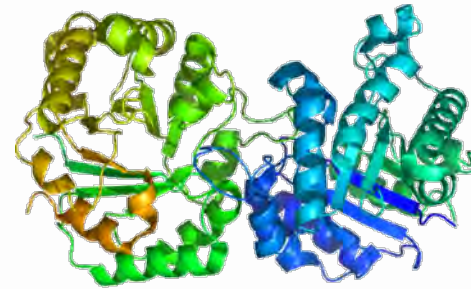
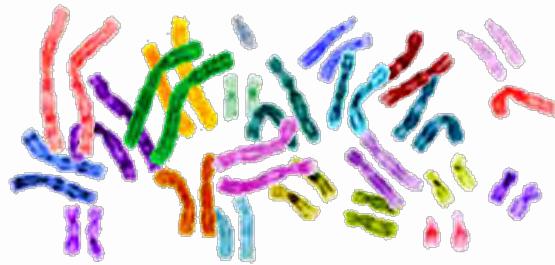


IBM

Key Innovation: Samples AND Handling Matter!

Genomics

Proteomics



All Depend On High-Quality,
Annotated Human Biospecimens

“Garbage In...Garbage Out”

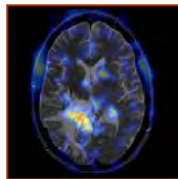
TCGA: Connecting Multiple Standardized Sources, Experiments, and Data Types



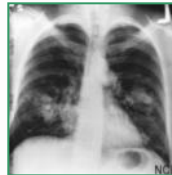
Three Cancers- Pilot

Multiple data types

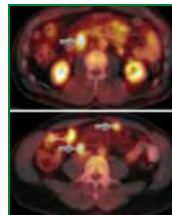
glioblastoma multiforme
(brain)



squamous carcinoma
(lung)



serous
cystadenocarcinoma
(ovarian)



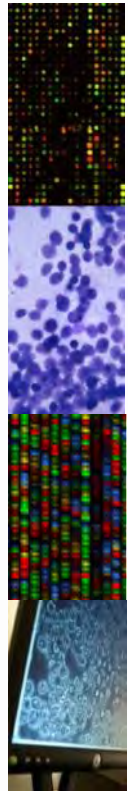
Biospecimen Core
Resource with more
than 13 Tissue
Source Sites

7 Cancer Genomic
Characterization
Centers

3 Genome
Sequencing
Centers

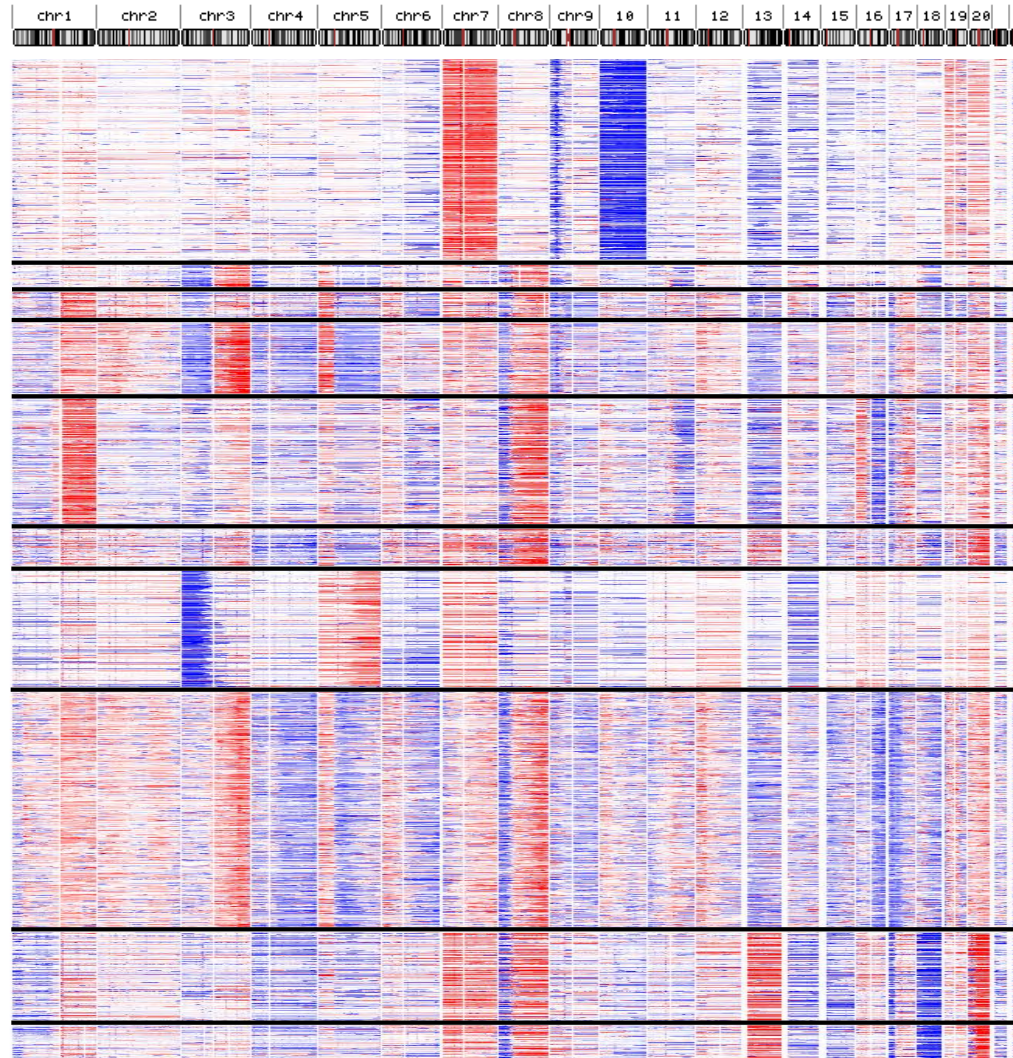
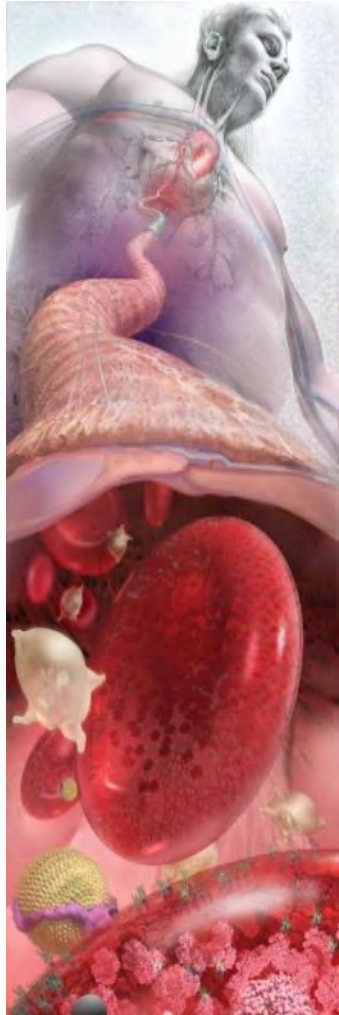
Data Coordinating
Center

- Clinical diagnosis
- Treatment history
- Histologic diagnosis
- Pathologic status
- Tissue anatomic site
- Surgical history
- Gene expression
- Chromosomal copy number
- Loss of heterozygosity
- Methylation patterns
- miRNA expression
- DNA sequence



Genomic “Steam Table”

Summer 2011



Glioblastoma:	470
Head & neck:	51
Lung adeno:	57
Lung squamous:	159
Breast carcinoma:	180
Stomach adeno:	84
Kidney clear carc:	260
Ovarian serous:	520
Colon adeno:	198
Rectum carcinoma:	

Total:

Genomic “Steam Table”

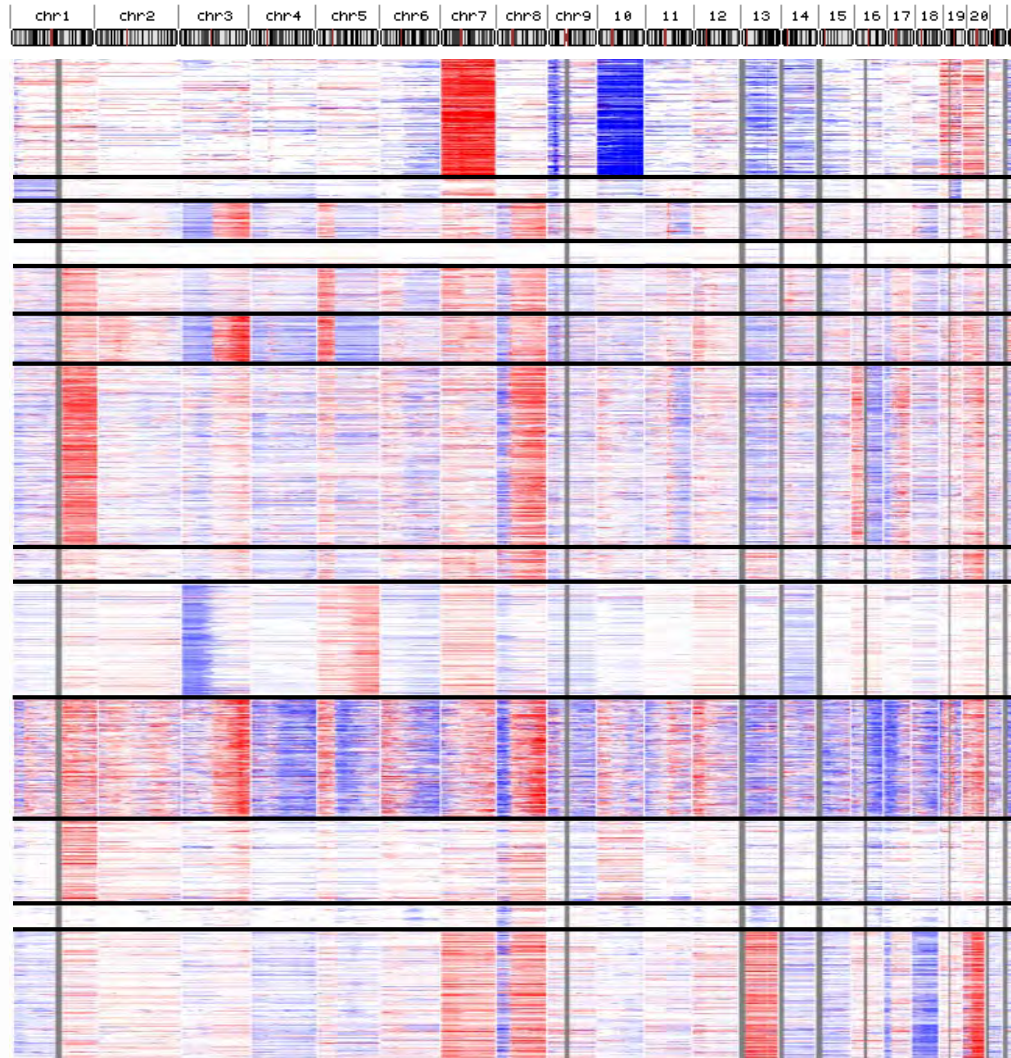
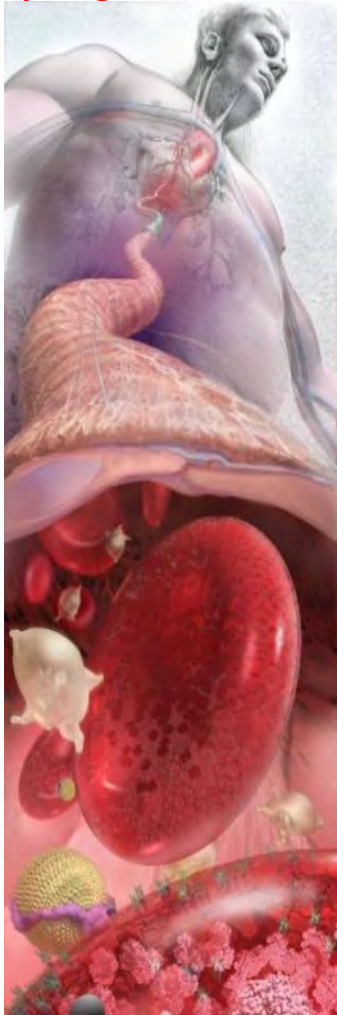
The Cancer Genome Atlas
Data Portal



Understanding genomics
to improve cancer care

CENTER for
STRATEGIC
SCIENTIFIC INITIATIVES

Spring 2012



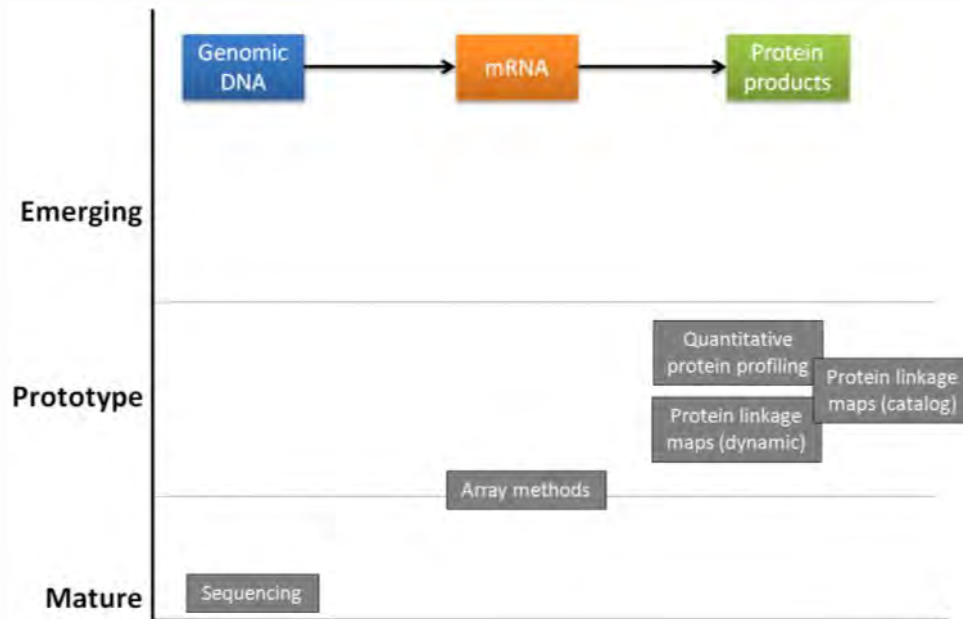
Glioblastoma:	535
Brain lower grade glioma:	80
Head & neck:	165
Thyroid carcinoma:	85
Lung adeno:	205
Lung squamous:	211
Breast carcinoma:	783
Stomach adeno:	149
Kidney clear carc:	489
Ovarian serous:	520
Uterine corpus end. car.:	363
Prostate adenocarcinoma:	82
Colon/rectum adeno:	564

Total:

Example #2: Cancer Proteomics and Mass Spectrometry

THE NATIONAL ACADEMIES
Advisers to the Nation on Science, Engineering, and Medicine

Technologies for Quantitative Analysis



Major Challenges

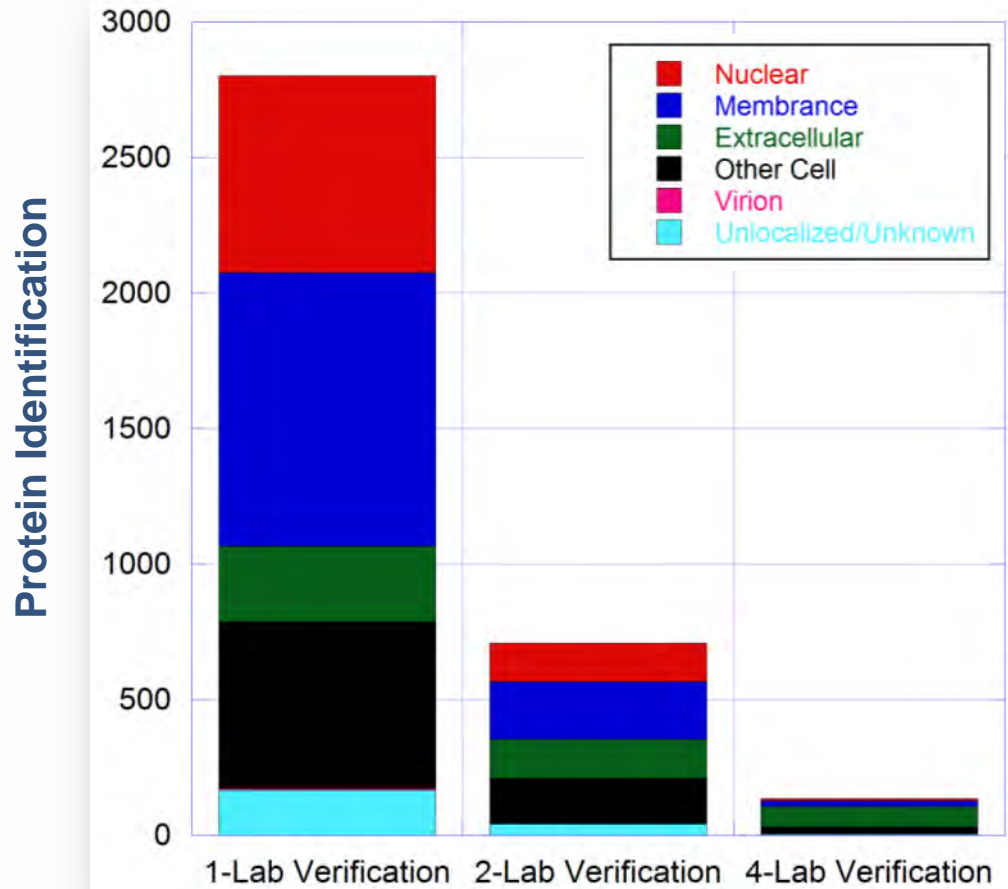
- Analytical variability in platforms
- Lack of standards, protocols, and reference data
- No consensus on data acquisition, analysis, and open access reporting of raw data

Unlike genomic technologies, proteomic technologies were not yet fully mature

Heterogeneity of Platforms and Reproducibility Challenges



Reproducibility of Clinical Proteomics in 2005



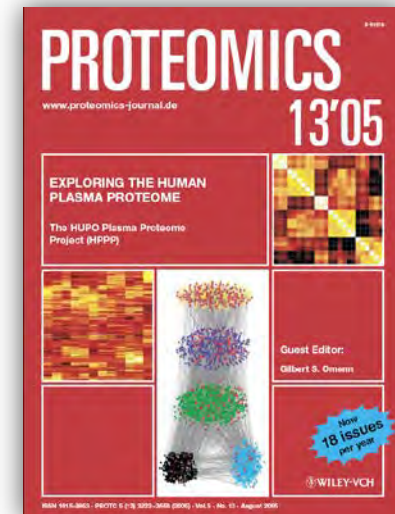
3226 DOI 10.1002/pmic.200500358 Proteomics 2005, 5, 3226-3245

REGULAR ARTICLE

2005

Overview of the HUPO Plasma Proteome Project: Results from the pilot phase with 35 collaborating laboratories and multiple analytical groups, generating a core dataset of 3020 proteins and a publicly-available database

Gilbert S. Omenn¹, David J. States¹, Marc Adamski¹, Thomas W. Blackwell¹, Rajasree Menon¹, Henning Hermjakob², Rolf Apweiler², Brian B. Haab³, Richard J. Simpson⁴, James S. Eddes⁴, Eugene A. Kapp⁴, Robert L. Moritz⁴, Daniel W. Chan⁵, Alex J. Rafi⁶, Arie Admon⁶, Ruedi Aebersold⁷⁻⁹, Jimmy Eng⁸, William S. Hancock⁸, Stanley A. Hefta¹⁰, Helmut Meyer¹¹, Young-Ki Paik¹², Jong-Shin Yoo¹³, Peipei Ping¹⁴, Joel Pounds¹⁵, Joshua Adkins¹⁵, Xiaohong Qian¹⁶, Rong Wang¹⁷, Valerie Wasinger¹⁶, Chi Yue Wu¹⁹, Xiaohang Zhao²⁰, Rong Zeng²¹, Alexander Archakov²², Akira Tsugita²³, Ilan Beer²⁴, Akhilesh Pandey⁵, Michael Pisano²⁵, Philip Andrews¹, Harald Tammen²⁶, David W. Speicher²⁷ and Samir M. Hanash^{1, 28}



Clinical Proteomic Technologies for Cancer (CPTAC) Pilot



nature
biotechnology

Multi-site assessment of the precision and reproducibility of multiple reaction monitoring-based measurements of proteins in plasma



Reproducibility

- **First demonstration** that MRM is highly reproducible **across multiple laboratories** and technology platforms
- **Community Resource:** Antibody Characterization Laboratory Launched

Data Sharing (“Amsterdam Principles”)

▪ Timing

- Data generated by individual investigators should **be released into the public domain at the latest upon publication while data generated by community resource projects should be released upon generation** following appropriate QA/QC procedures

▪ Comprehensiveness

- High quality raw data (e.g., mass spectral, protein/affinity array data) be released to the public. They should be well annotated with metadata, information on data quality, and identification quality control data

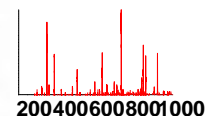
▪ Format

- Open access to proteomic data requires community-supported standardized formats, controlled vocabularies, reasonable reporting requirements, and publicly available central repositories

research articles **proteome** May 2009
**Recommendations from the 2008 International Meeting on Proteomics
Data Release and Sharing Policy: The Amsterdam Principles**

Henry Rodriguez,^{1,2} Mike Snyder,³ Mathias Uhlen,⁴ Phil Andrews,⁵ Ronald Beavis,⁶ Christoph Borchers,⁷ Robert J. Chalkley,⁸ Sung Yun Cho,⁹ Katie Cottingham,¹⁰ Michael Dunn,¹¹ Tomasz Dylag,¹² Ron Edgar,¹³ Peter Hare,¹⁴ Albert J. R. Heck,¹⁵ Roland F. Hirsch,¹⁶ Karen Kennedy,¹⁷ Patrik Kolar,¹⁸ Hans-Joachim Kraus,¹⁹ Parag Mallick,²⁰ Alexey Nesvizhskii,²¹ Peipei Ping,²² Fredrik Pontén,²³ Liming Yang,²⁴ John R. Yates,²⁵ Stephen E. Stein,²⁶ Henning Hermjakob,²⁷ Christopher R. Kinsinger,²⁸ and Rolf Apweiler²⁹

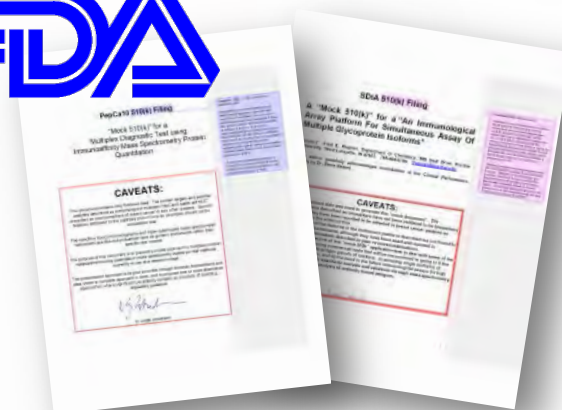
Center for Strategic Scientific Initiatives, National Cancer Institute, National Institutes of Health, Bethesda, Maryland, 20832, Department of Molecular, Cellular, and Developmental Biology, Yale University, New Haven, Connecticut 06520, KTH Biotechnology, KTH - AlvaNova University Center, Stockholm, Sweden, Department of Biological Chemistry, University of Michigan Medical School, Ann Arbor, Michigan 48109, Department of Medical Genetics, University of British Columbia, Vancouver, British Columbia, Canada, University of Victoria Proteomics Centre, Victoria, British Columbia, Canada, Department of Pharmaceutical Chemistry, University of California, San Francisco, San Francisco, California, 94158, Yonsei Proteome Research Center, Yonsei



CPTAC Pilot: Crosstalk with FDA and Educating Community

- **Analytical Validation Review Documents**

- CPTAC/FDA Workshop - identify analytical validation needs for clinical proteomic technologies in the context of intended use



- **Outputs:**

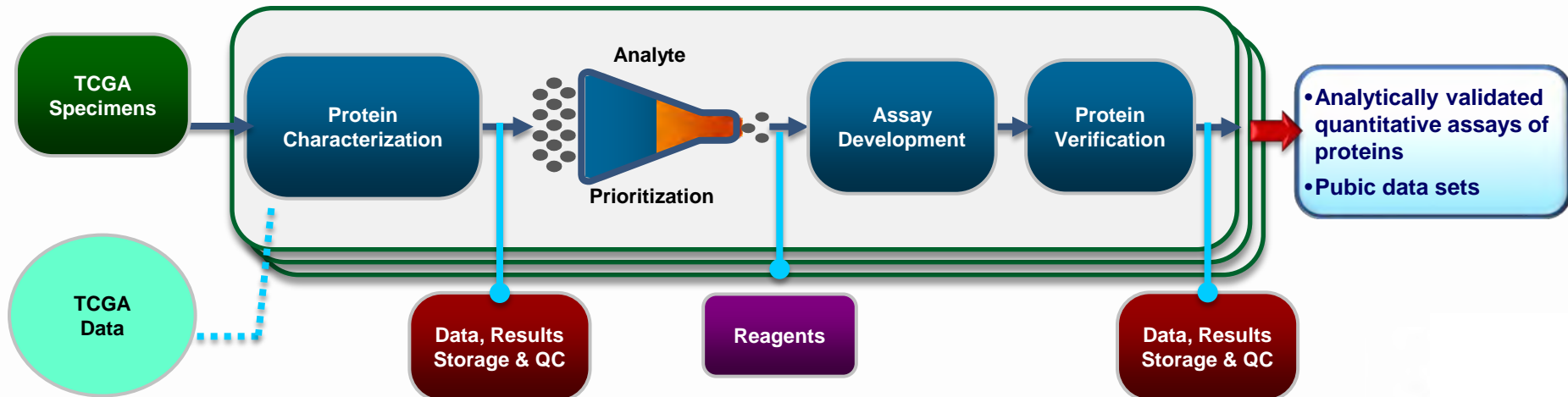
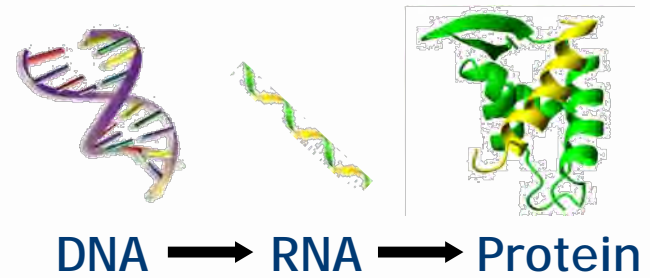
- Public mock 510(k) pre-applications that serve as review documents on:
 - multiplex MRM-MS assay
 - multiplex affinity-based assay
- Published in special issue of *Clinical Chemistry* (by AACC), that **informs research community and FDA to technology platforms that will likely be part of future 510k submissions**



Clinical Proteomic Tumor Analysis Centers (CPTAC Phase II)

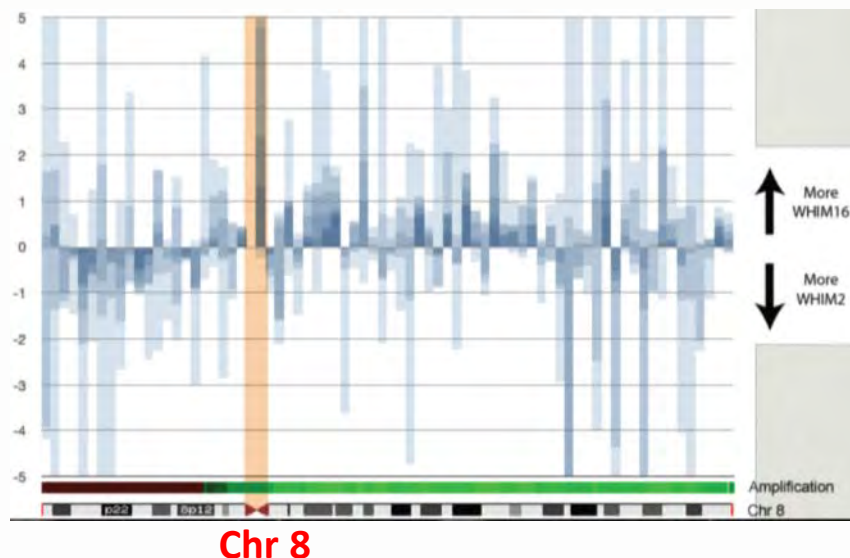
Phase II Launched Sept 2011

- Analyze matched TCGA samples using two approaches
 - Targeting genome to proteome
 - Mapping proteome to genome
- Develop validated and quantitative assays and reagents
- Distribute raw and analyzed data via public data portal

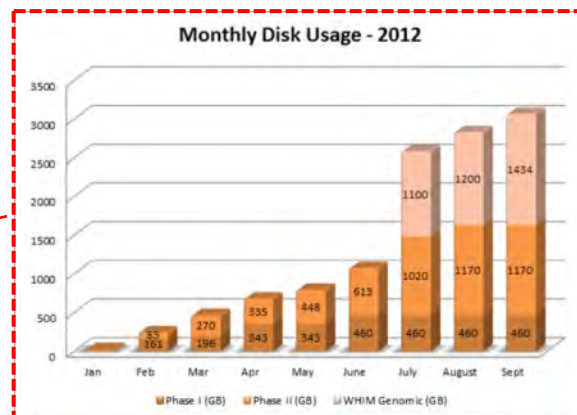


CPTAC Phase II: Highlights, Progress to Date, and Data Release

Status Update: Fall 2012

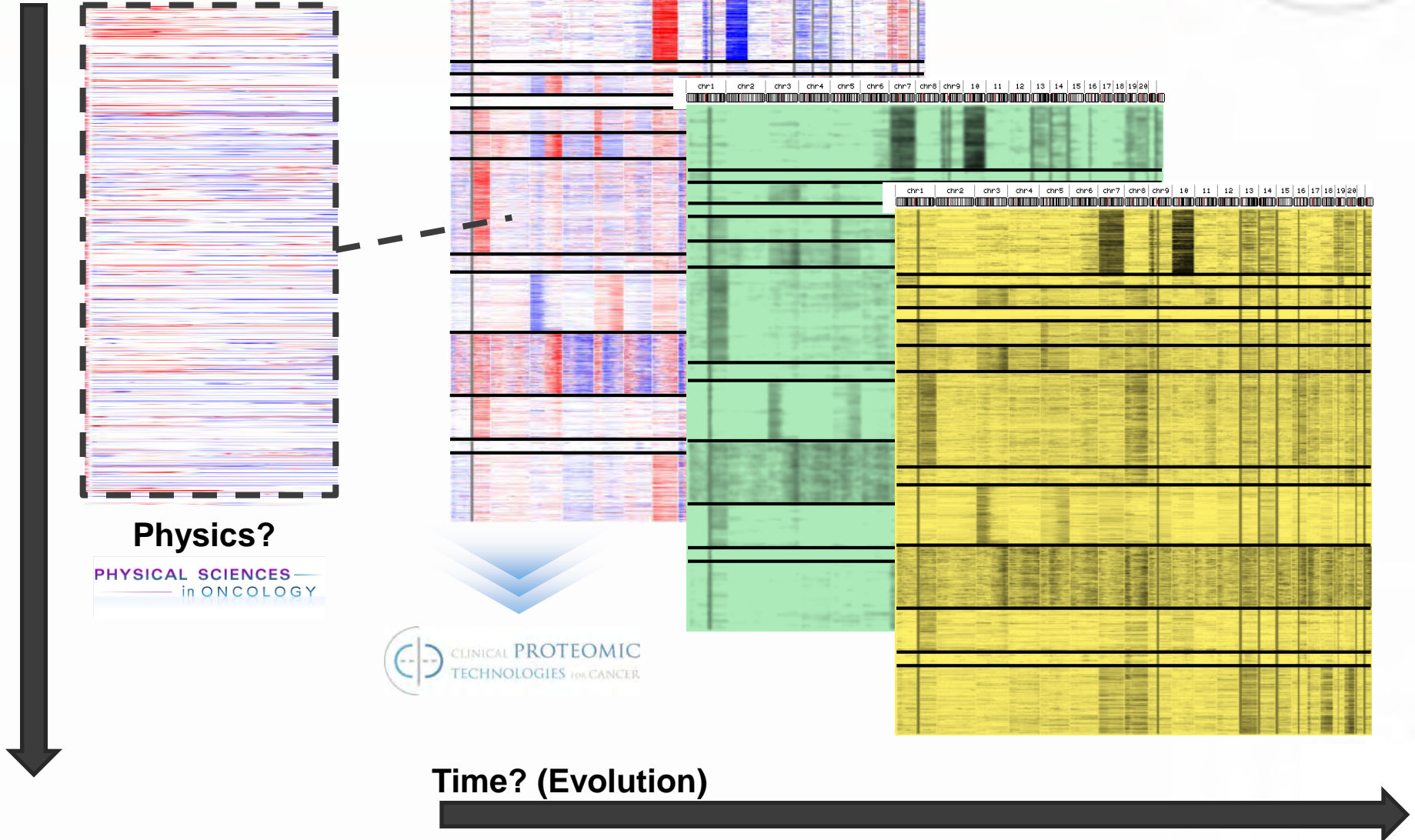


- **Due diligence studies** near completion
 - Cross network experiments show comparable lab-to-lab measurements
- **Orthogonal** proteomic platforms and analysis (proteome → genome vs. genome → proteome) reveal **additional unexpected complexities**
- Verifying new insights will require **additional sample sets** and **development of novel analysis** algorithms and techniques
- Public data portal access **OPEN!**
(Phase I: 351 GB, Phase II: 616 GB)



Where Do We Go From Here?

Is it JUST More Data?



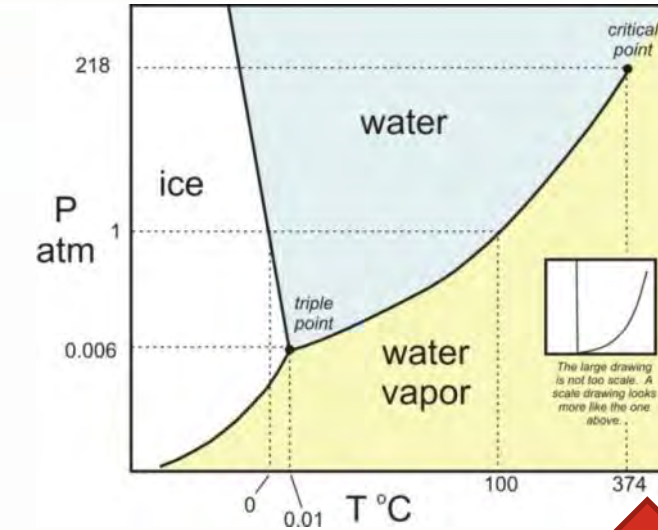
Standards and Sharing of Data → New Insights and Understanding

- Define samples & protocols
 - Share collected data



2400m

0m



New Understanding

- Phase boundaries
 - V/L equilibrium
 - Triple Point**

(Phase Diagram)

New Parameter

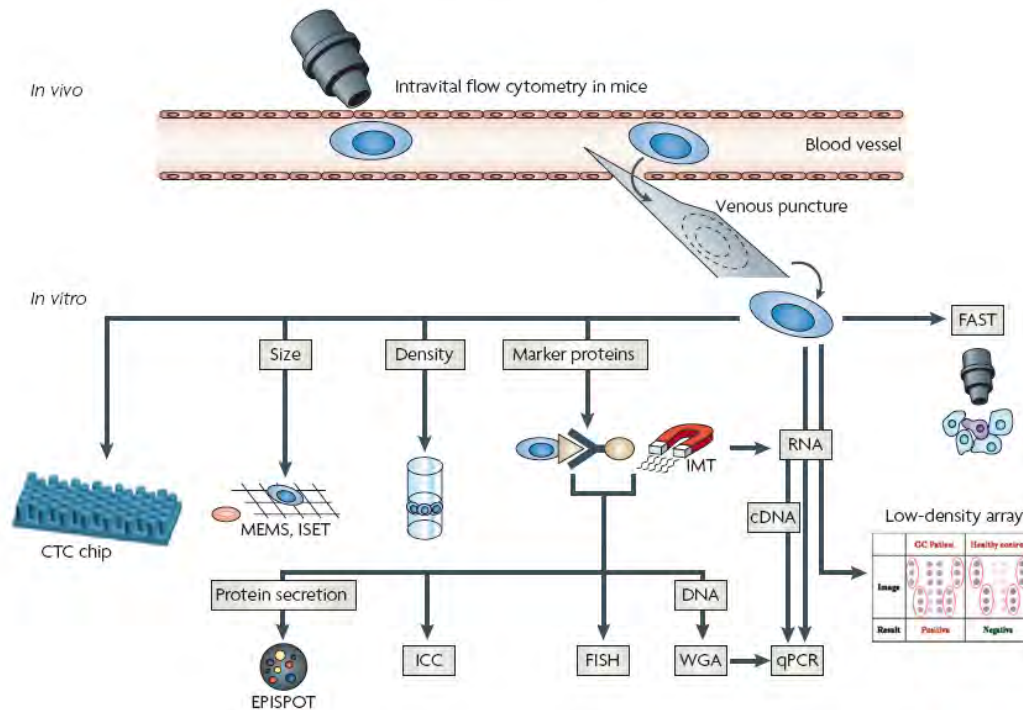
"Pressure"

Pressure (kg/cm ²)	Temp (°C)	Vapour enthalpy (kcal/kg)	Specific volume (m ³ /kg)	Superheated steam	
				Density (kg/m ³)	Specific volume (m ³ /kg) at 250°C at 300°C
1	99.1	638.8	1.725	0.580	2.454 2.691
2	119.6	646.2	0.902	1.109	1.223 1.342
3	132.9	650.6	0.617	1.621	0.812 0.893
4	142.9	653.7	0.471	2.123	0.607 0.668
5	151.1	656.0	0.382	2.618	0.484 0.533
6	158.1	657.0	0.321	3.115	0.402 0.443
7	164.2	659.5	0.278	3.597	0.343 0.379
8	169.6	660.8	0.245	4.082	0.299 0.331
9	174.5	661.9	0.219	4.566	0.265 0.293
10	179.1	662.9	0.198	5.051	0.238 0.263
12	187.1	664.5	0.166	6.024	0.196 0.218
14	194.1	665.7	0.143	6.993	0.167 0.186
16	200.4	666.7	0.126	7.937	0.145 0.162
18	206.1	667.4	0.112	8.929	0.128 0.143
20	211.4	668.0	0.101	9.901	0.114 0.128
22	216.2	668.4	0.092	10.870	0.103 0.116
24	220.7	668.7	0.085	11.765	0.093 0.106
26	225.0	669.0	0.078	12.821	0.085 0.097
28	229.0	669.1	0.073	13.699	0.078 0.089
30	232.7	669.2	0.068	14.706	0.072 0.083

LOTS of
Quantitative
and
Reproducible
Data

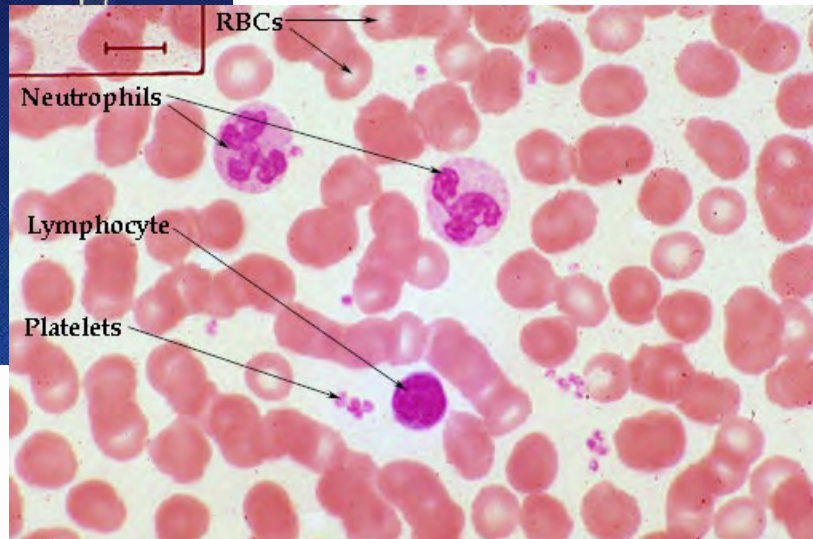
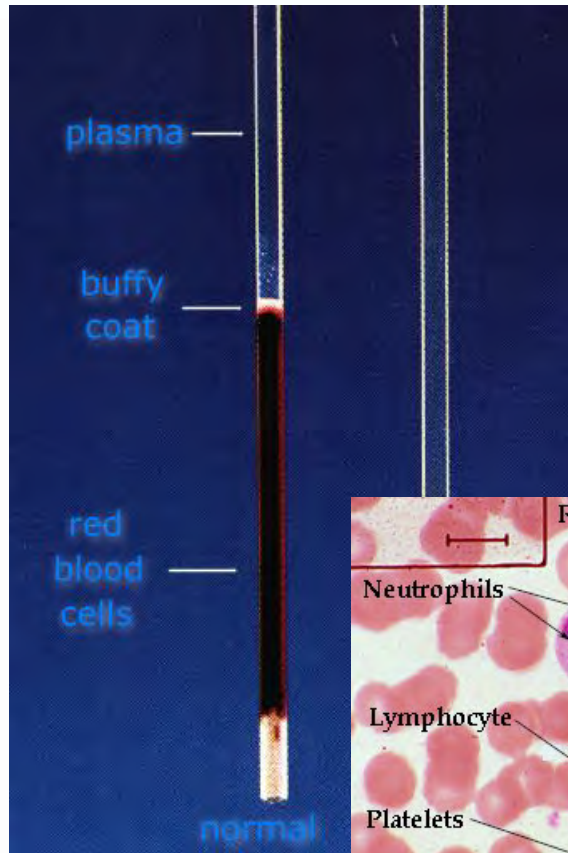
(Steam Table)

Your Dilemma: Finding the “needle”?



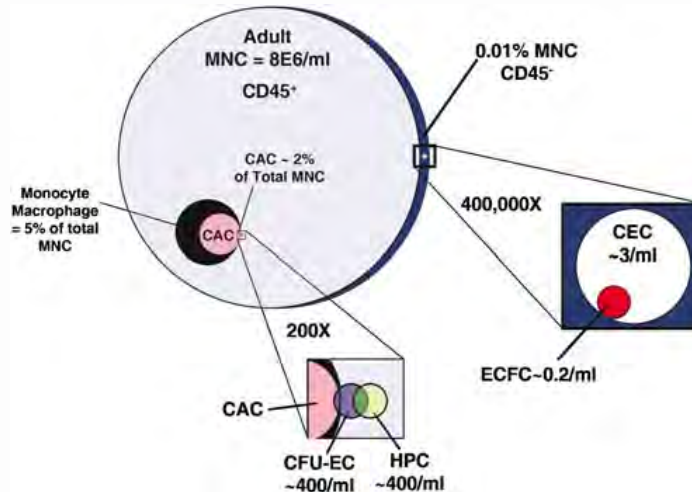
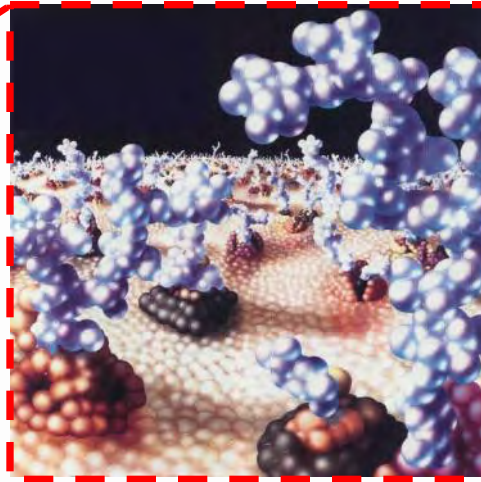
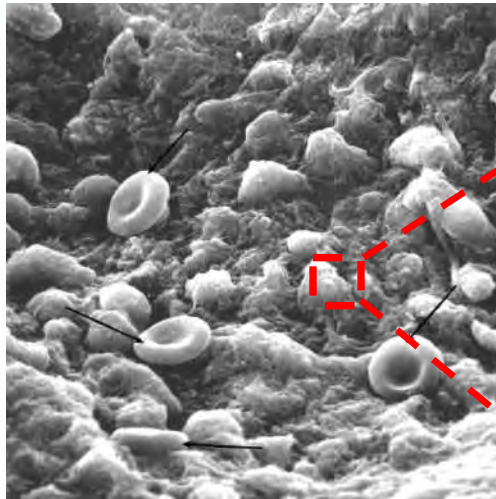
- Understanding of tumor cells in the blood has gained considerable attention in recent years, mainly due to advancements in technology
 - Disseminated tumor cells (DTCs)** in the bone marrow
 - Circulating tumor cells (CTCs)** in peripheral blood
- ~ 1 DTC/CTC per $10^6 - 10^7$ nucleated blood or BM cells

The “hay”: the many components of blood



- ~5 liters
- Major elements
 - ~**55% Plasma**
 - Mostly water
 - Serum- clotting agents removed
 - ~**45% erythrocytes**
 - ~4-6 billion/ml
 - ~2-3 million/second
 - ~**1% leukocytes**
 - ~6-10 million/ml
 - ~**thrombocytes**
 - ~200-500 million/ml

Sorting out which “needle”



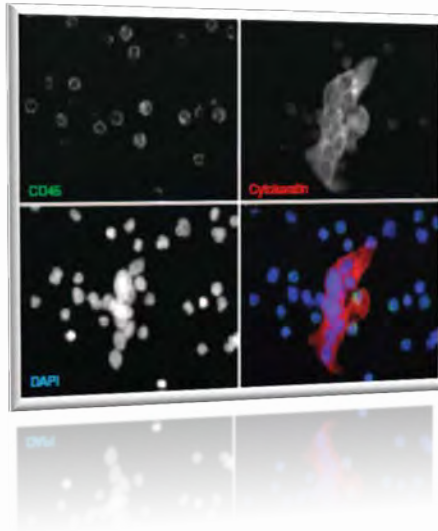
- Finding A “needle” is hard enough
- Even harder if there are “decoys” hidden within

Timing Seems Right...

NEWS FEATURE

Beyond counting tumor cells

Since the discovery of circulating tumor cells in 1869, researchers have been able to do little else beyond count them. This is about to change, as advanced technologies for harvesting and analyzing rare cells from blood are opening the window for characterization. Jim Kling reports.



REVIEW

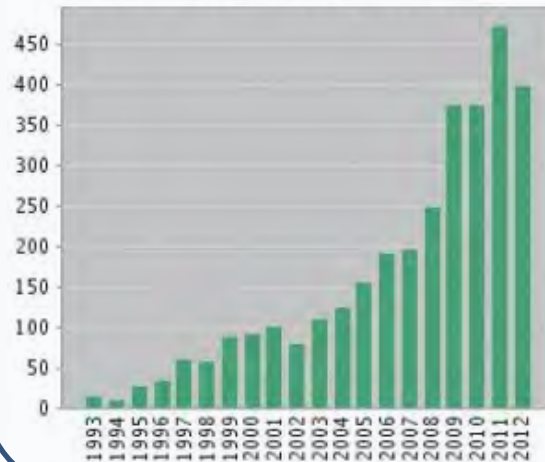


JOURNAL OF
TRANSLATIONAL MEDICINE

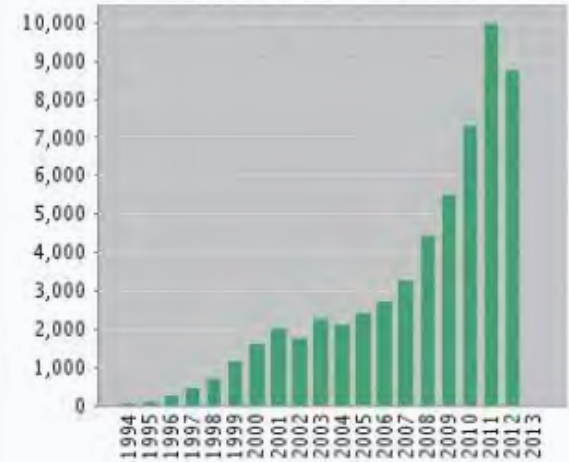
Considerations in the development of circulating tumor cell technology for clinical use

David R Parkinson^{1*}, Nicholas Dracopoli², Brenda Gumbs Petty³, Carolyn Compton⁴, Massimo Cristofanilli⁵, Albert Deisseroth⁶, Daniel F Hayes⁷, Gordon Kapke⁸, Prasanna Kumar⁹, Jerry SH Lee¹⁰, Minetta C Liu¹¹, Robert McCormack¹², Stanislaw Mikulski¹³, Larry Nagahara¹⁰, Klaus Pantel¹⁴, Sonia Pearson-White¹⁵, Elizabeth A Punnoose¹⁶, Lori T Roadcap¹⁷, Andrew E Schade¹⁸, Howard I Scher¹⁹, Caroline C Sigman³ and Gary J Kelloff¹⁰

Published Items in Each Year



Citations in Each Year



Scripps PSOC Clinical Studies

LUNG:

- **PSOC0043** (UCSD, Billings)
- **PSOC0044** (Scripps Clinic, UCSD, Billings)
- **PSOC0046** (UCSD)
- **PSOC0047** (NKI, UCSD, Billings)
- **PSOC0048** (NKI, Amsterdam)
- **PSOC0049** (Stanford, USC)
- **PSOC0064** (UCSD)
- **PSOC0065** (UCSD)

LIVER:

- **PSOC0050** (Scripps Green Hospital, Scripps Clinic)
- **PSOC0055** (UCSF, Northwestern University)

SKIN:

- **PSOC0056** (Pacific Oncology and Hematology)
- **PSOC0061** (Nevada Cancer Center)

PROSTATE:

- **PSOC0045** (Scripps Clinic: Anderson Outpatient Pavillion, Carmel Valley, Rancho Bernardo, Torrey Pines Science Park)
- **PSOC0051** (USC)
- **PSOC0058** (USC)
- **PSOC0060** (Scripps Health)
- **PSOC0063** (NorthShore)

BREAST:

- **PSOC0045** (Scripps Clinic: Anderson Outpatient Pavillion, Carmel Valley, Rancho Bernardo, Torrey Pines Science Park)
- **PSOC0053** (Duke University)
- **PSOC0060** (Scripps Health)
- **PSOC0062** (Scripps Clinic: Anderson Outpatient Pavillion, Carmel Valley, Rancho Bernardo, Torrey Pines Science Park)

HEART:

- **PSOC0057** (Scripps Health)

PANCREAS:

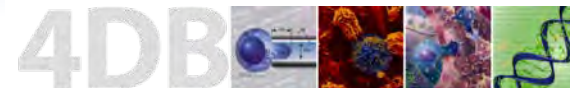
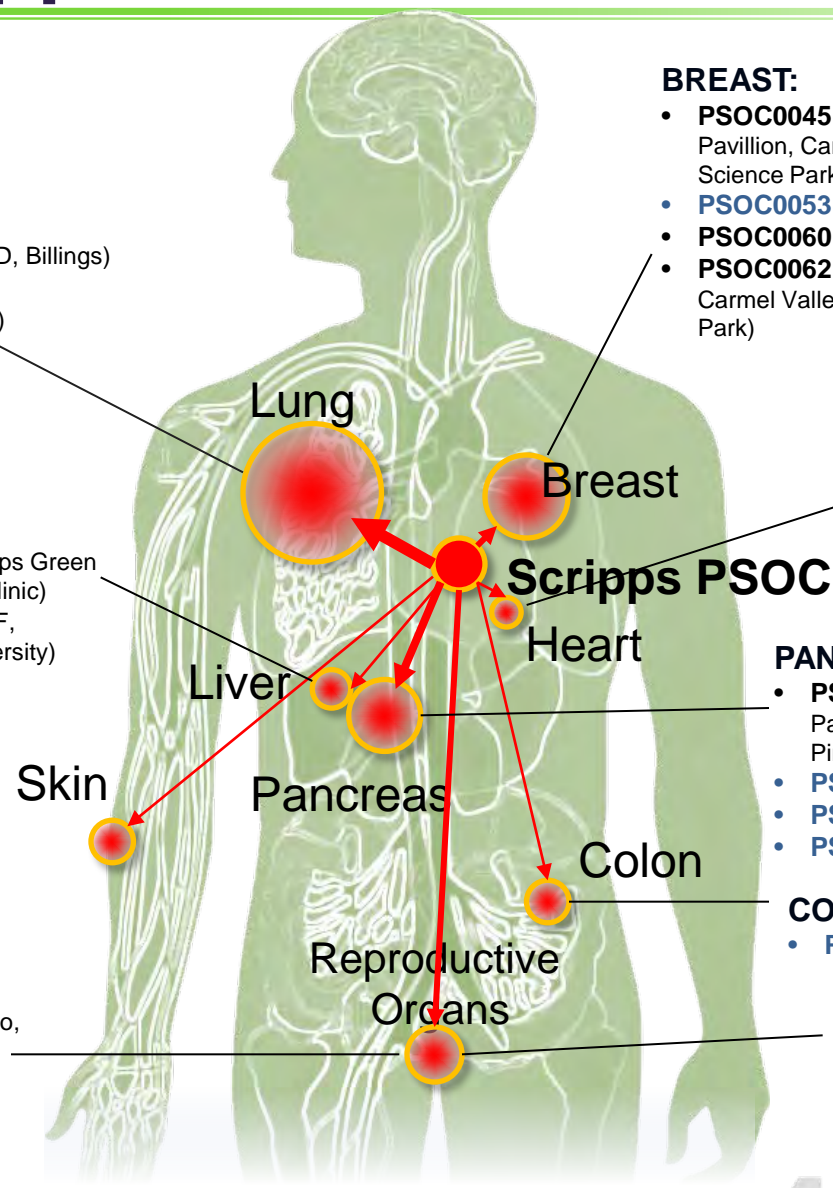
- **PSOC0045** (Scripps Clinic: Anderson Outpatient Pavillion, Carmel Valley, Rancho Bernardo, Torrey Pines Science Park)
- **PSOC0054** (UCSF)
- **PSOC0059** (Scripps Green Hospital)
- **PSOC0060** (Scripps Health)

COLON:

- **PSOC0066** (Scripps Clinic)

OVARIAN:

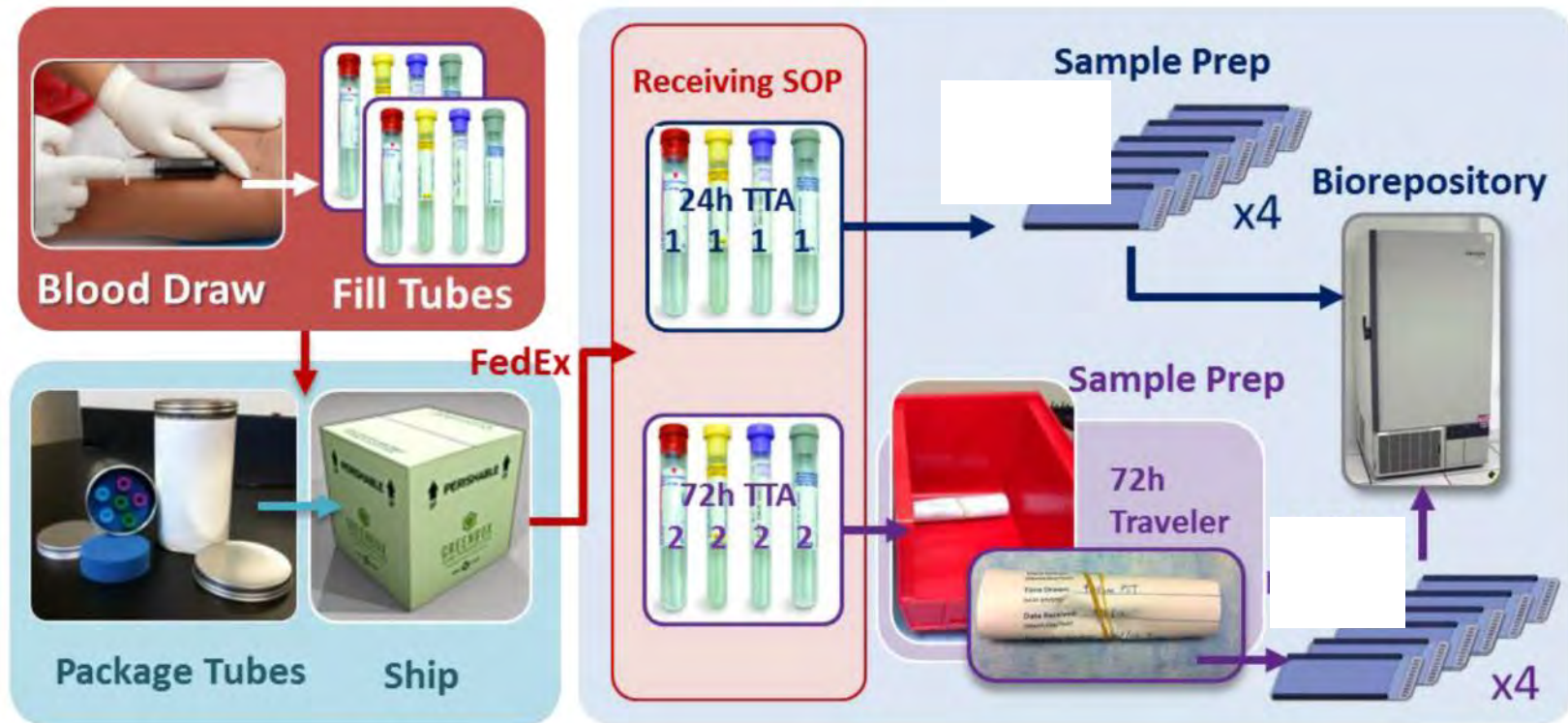
- **PSOC0052** (Scripps Memorial Hospital, South Coast Gynecologic Oncology)



Pilot Project: Impact of pre-analytical variables on CTCs

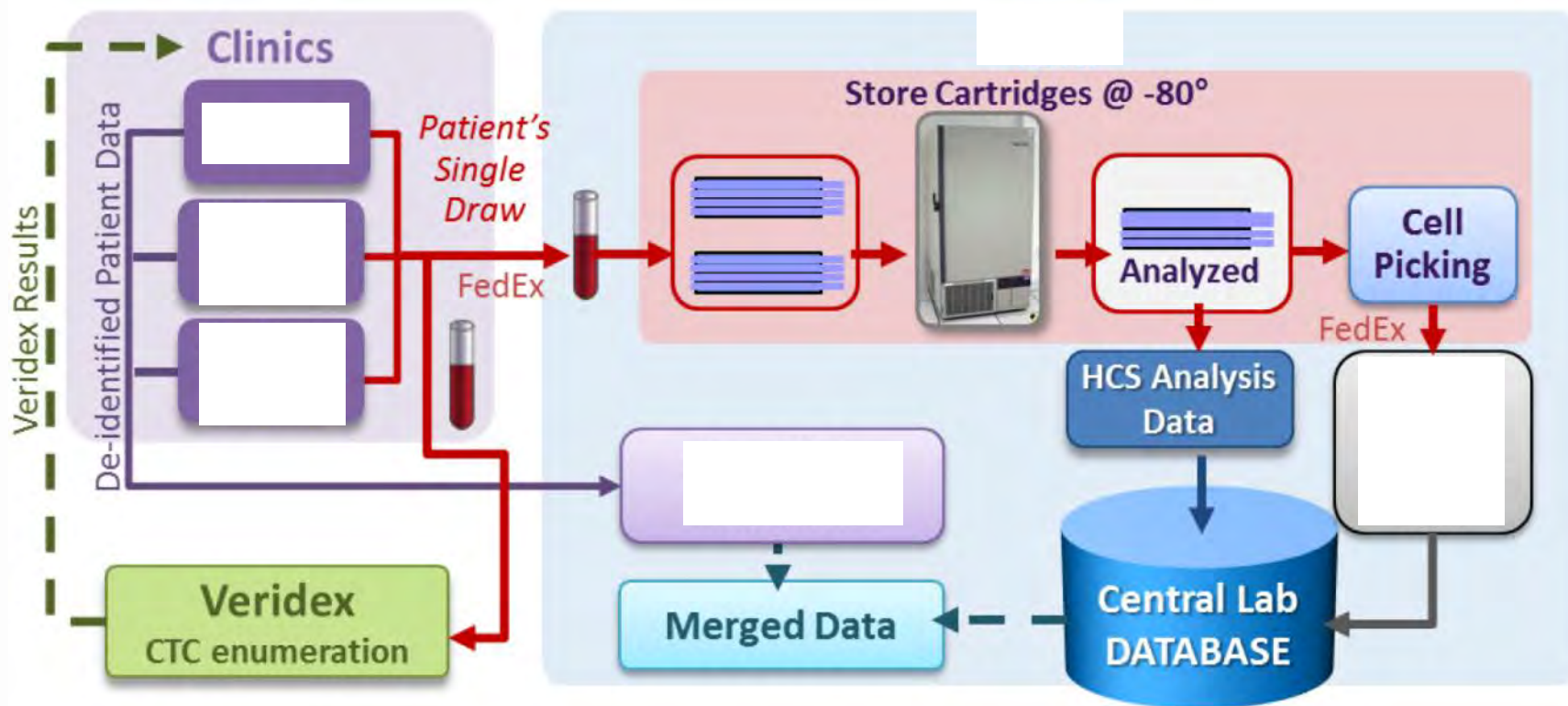
Project Objective:

- Strengthen the research use and clinical utility of HCS CTC assays
- Develop SOPs using the best pre-analytical conditions for blood CHP (collection, handling, processing)
 - Blood collection tube type (Streck, EDTA, Citrate, Heparin)
 - Time to assay (24 and 72 hours)

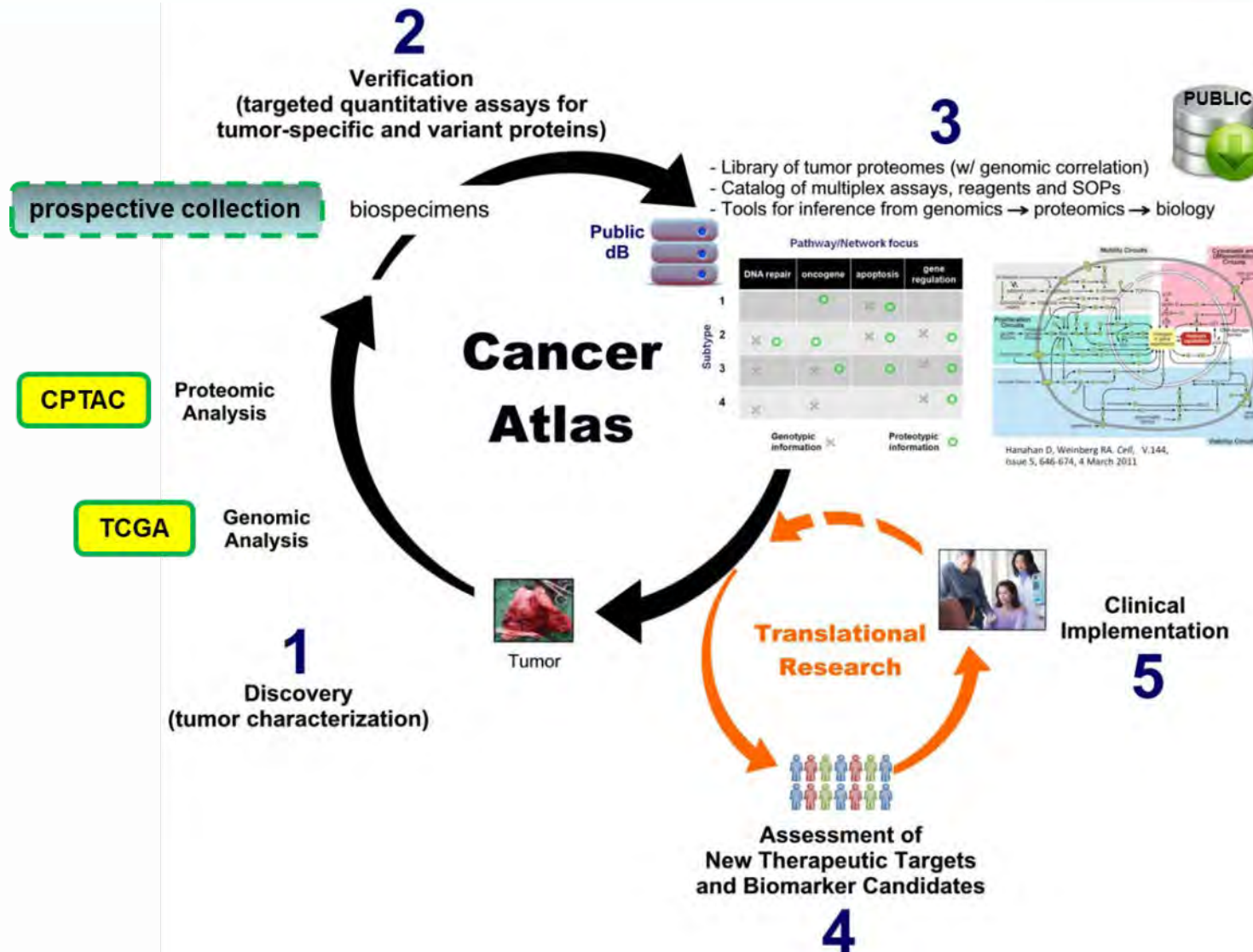


Key Pilot Output: Public Release of Data

Results will be compared to the Veridex CellSearch® system and
will be deposited in a publically available database



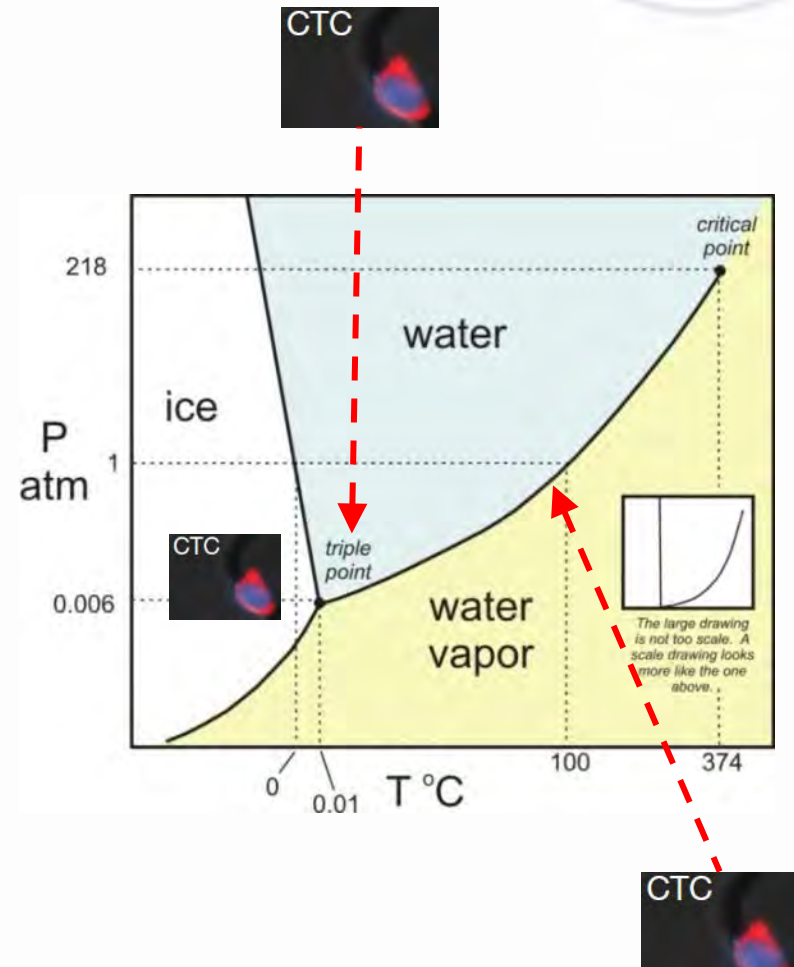
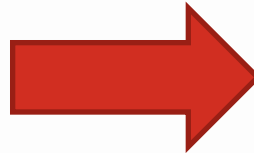
Is There A Place for CTCs...



Personal Prediction...

**“LOTS of
Quantitative and
Reproducible
Data”**

Saturated steam				Superheated steam		
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28	229.0	669.1	0.073	13.699	0.078	0.089
30	232.7	669.2	0.068	14.706	0.072	0.083



Relevant CSSI Funding Opportunities



http://cssi.cancer.gov/resources-current_funding.asp

Provocative Questions (\$30M)

Due Date 12/04/12

- **Research Answers to NCIs Provocative Questions**
 - **Group A:** RFA-CA-12-015 (R01) [\$5-\$7M]
RFA-CA-12-016 (R21) [\$2-\$3M]
 - **Group B:** RFA-CA-12-017 (R01) [\$5-\$7M]
RFA-CA-12-018 (R21) [\$2-\$3M]
 - **Group C:** RFA-CA-12-019 (R01) [\$5-\$7M]
RFA-CA-12-020 (R21) [\$2-\$3M]
 - **Group D:** RFA-CA-12-021 (R01) [\$5-\$7M]
RFA-CA-12-022 (R21) [\$2-\$3M]

Innovative Molecular Analysis Technologies (\$10.5M)

Due Date 02/20/13

- **Early-Stage Innovative Technology Development (R21)**
 - RFA-CA-13-001 (R21, 3 years) [\$5M]
- **Validation and Advanced Development of Emerging Technologies (R33)**
 - RFA-CA-13-002 (R33) [\$3.5M]
- **Early-Stage and/or Validating Technologies in Biospecimen Science (R21/R33)**
 - RFA-CA-13-003 (R21) [\$0.8M]
 - RFA-CA-13-004 (R33) [\$0.7M]



Learn More About CSSI/CCG...



<http://cssi.cancer.gov>



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IMAT Program Director

Tony Dickherber, PhD



Jerry S.H. Lee, PhD, jerry.lee@nih.gov